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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/543,003	04/24/2006	Yuichi Ono	082368-004900US	7150
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TWO EMBAR	CADERO CENTER	CKLW, LLI	LANDSMAN	, ROBERT S
EIGHTH FLO SAN FRANCI	OR SCO, CA 94111-3834		ART UNIT	PAPER NUMBER
	,		1647	
		•	MAIL DATE	DELIVERY MODE
			05/15/2007	PAPER

Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

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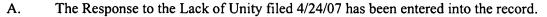
		Application N	o. Applica	nt(s)
•		10/543,003	ONO ET	· AL.
	Office Action Summary	Examiner	Art Unit	
		Robert Landsn	nan, Ph.D. 1647	
Period fo	The MAILING DATE of this communica or Reply	tion appears on the cov	er sheet with the correspon	dence address
WHIC - Exter after - If NO - Failu Any r	ORTENED STATUTORY PERIOD FOR CHEVER IS LONGER, FROM THE MAIL asions of time may be available under the provisions of 3 SIX (6) MONTHS from the mailing date of this communic period for reply is specified above, the maximum statute to reply within the set or extended period for reply will, eply received by the Office later than three months after ad patent term adjustment. See 37 CFR 1.704(b).	LING DATE OF THIS (7 CFR 1.136(a). In no event, ho cation. ary period will apply and will expi by statute, cause the application	COMMUNICATION. Inwever, may a reply be timely filed The SIX (6) MONTHS from the mailing of the to become ABANDONED (35 U.S.C.)	date of this communication.
Status	\			
1)⊠	Responsive to communication(s) filed of	on 24 April 2007.		
•	•	☐ This action is non-f	nal.	
′=	Since this application is in condition for			as to the merits is
7,—	closed in accordance with the practice	•	•	
Dispositi	on of Claims			
4)⊠	Claim(s) 1.3 and 5-9 is/are pending in t	he application.	•	
•	4a) Of the above claim(s) <u>3 and 5-9</u> is/a		sideration.	
5)	Claim(s) is/are allowed.			
6)🖾	Claim(s) 1 is/are rejected.	·		
7)	Claim(s) is/are objected to.		•	
8)[Claim(s) are subject to restriction	n and/or election requi	ement.	
Applicati	on Papers			
9)🛛	The specification is objected to by the E	xaminer.		
10)⊠	The drawing(s) filed on <u>22 July 2005</u> is/s	are: a)⊠ accepted or	b) objected to by the Ex	aminer.
	Applicant may not request that any objectio	n to the drawing(s) be he	ld in abeyance. See 37 CFR	1.85(a).
	Replacement drawing sheet(s) including the	e correction is required if	the drawing(s) is objected to.	See 37 CFR 1.121(d).
11)	The oath or declaration is objected to by	y the Examiner. Note the	ne attached Office Action of	r form PTO-152.
Priority u	ınder 35 U.S.C. § 119			
a)[Acknowledgment is made of a claim for All b) Some * c) None of: 1. Certified copies of the priority do: 2. Certified copies of the priority do: 3. Copies of the certified copies of the application from the International see the attached detailed Office action for	cuments have been recuments have been reche priority documents Bureau (PCT Rule 17	ceived. ceived in Application No have been received in this .2(a)).	
2) Notic 3) Inform	t(s) e of References Cited (PTO-892) e of Draftsperson's Patent Drawing Review (PTO nation Disclosure Statement(s) (PTO/SB/08) r No(s)/Mail Date <u>7/22/05</u> .	-948) 5) [☐ Interview Summary (PTO-413) Paper No(s)/Mail Date. ☐ Notice of Informal Patent Applic ☐ Other: <u>Sequence Comparison</u> .	

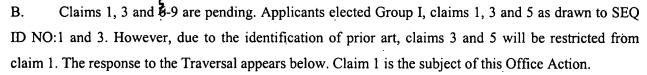
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DETAILED ACTION

1. Formal Matters





2. Rejoinder

A. The examiner has required restriction between product and process claims. Where applicant elects claims directed to the product, and a product claim is subsequently found allowable, withdrawn process claims that depend from or otherwise include all the limitations of the allowable product claim will be rejoined in accordance with the provisions of MPEP § 821.04. Process claims that depend from or otherwise include all the limitations of the patentable product will be entered as a matter of right if the amendment is presented prior to final rejection or allowance, whichever is earlier. Amendments submitted after final rejection are governed by 37 CFR 1.116; amendments submitted after allowance are governed by 37 CFR 1.312.

In the event of rejoinder, the requirement for restriction between the product claims and the rejoined process claims will be withdrawn, and the rejoined process claims will be fully examined for patentability in accordance with 37 CFR 1.104. Thus, to be allowable, the rejoined claims must meet all criteria for patentability including the requirements of 35 U.S.C. 101, 102, 103, and 112. Until an elected product claim is found allowable, an otherwise proper restriction requirement between product claims and process claims may be maintained. Withdrawn process claims that are not commensurate in scope with an allowed product claim will not be rejoined. See "Guidance on Treatment of Product and Process Claims in light of *In re Ochiai, In re Brouwer* and 35 U.S.C. § 103(b)," 1184 O.G. 86 (March 26, 1996). Additionally, in order to retain the right to rejoinder in accordance with the above policy, Applicant is advised that the process claims should be amended during prosecution either to maintain dependency on the product claims or to otherwise include the limitations of the product claims. Failure to do so may result in a loss of the right to rejoinder.

Further, note that the prohibition against double patenting rejections of 35 U.S.C. 121 does not apply where the restriction requirement is withdrawn by the examiner before the patent issues. See MPEP § 804.01.



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3. Traversal

A. Applicants argue that a special technical feature, the dopaminergic cell marker, exists among Groups I-IV. Regarding the traversal of SEQ ID NO:1 vs. 2 and 3 vs 4. Applicants have stated that these polynucleotides and polypeptides are 82-83% identical. These arguments have been considered, but is not deemed persuasive. First, as seen below, there is prior art under 35 USC 102 on the special technical feature (i.e. the marker). Furthermore, Applicants have not identified a common core structure between the proteins of SEQ ID NO:2 and 4, or the nucleic acids of SEQ ID NO:1 and 3. However, since SEQ ID NO:1 encodes SEQ ID NO:3, these two sequence will be searched. This Lack of Unity is deemed proper and is, therefore, made FINAL.

4. Specification

- A. The specification is objected to due to the use of hyperlinks, for example, on page 25, lines 1 and 5. Applicant is advised that embedded hyperlinks and/or other forms of browser-executable code are impermissible and require deletion. The attempt to incorporate subject matter into the patent application by reference to a hyperlink and/or other forms of browser-executable code is considered to be an improper incorporation by reference. See MPEP 608.01(p), paragraph I regarding incorporation by reference.
- B. The specification is objected to since the Brief Description of Figure 7 should begin with, e.g, "Fig. 7A-D are..."
- C. According to 37 CFR 1.821(d) (MPEP § 2422), where the description or claims of a patent application discuss a sequence listing that is set forth in the "Sequence Listing" in accordance with paragraph (c) of this section, reference must be made to the sequence by use of the assigned identifier, in the text of the description or claims, even if the sequence is also embedded in the text of the description or claims of the patent application. Sequences appear on page 33 lines 1-3 of the specification but are not identified by SEQ ID NO as required.

5. Claim Objections

A. Claim 1 is objected to for reciting non-elected subject matter (SEQ ID NO:2 and 4).

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B. No objection is being made to claim 1 regarding the phrase "a nucleotide sequence." However, it is brought to Applicants' attention that the use of the word "a" can be interpreted, for prior art purposes, as meaning a single nucleotide or amino acid – as opposed to "the nucleotide sequence" which is interpreted as the entire sequence.

6. Claim Rejections - 35 USC § 112, first paragraph - scope of enablement

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

A. Claims 1 is rejected under 35 U.S.C. 112, first paragraph, because the specification, while being enabling for the marker of SEQ ID NO:1, does not reasonably provide enablement for markers which are polynucleotides which are fragments of, or which hybridize to, SEQ ID NO:1. The specification does not enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make the invention commensurate in scope with these claims.

In <u>In re Wands</u>, 8USPQ2d, 1400 (CAFC 1988) page 1404, the factors to be considered in determining whether a disclosure would require undue experimentation include (1) the quantity of experimentation necessary, (2) the amount of direction or guidance presented, (3) the presence or absence of working examples, (4) the nature of the invention, (5) the state of the prior art, (6) the relative skill of those in the art, (7) the predictability or unpredictability of the art, and (8) the breadth of the claims.

First, the breadth of the claim is excessive with regard to claiming all polynucleotide markers which "hybridize" under stringent conditions to that of SEQ ID NO:1, or which are "fragments" of SEQ ID NO:1. Nucleic acid molecules which "hybridize" to SEQ ID NO:1 would have one or more nucleic acid substitutions, deletions, insertions and/or additions to said polynucleotides.

Applicants provide no guidance or working examples of nucleic acid molecules which hybridize to SEQ ID NO:1, or which fragments of SEQ ID NO:1 can be used as markers, nor is it predictable to one of ordinary skill in the art what polynucleotides which hybridize to SEQ ID NO:1 could be used as markers for the claimed cells. Furthermore, the polynucleotide of SEQ ID NO:1 is over 4000 bases. It is not known which fragments of comprising only 15 bases could be used to identify the claimed cell. Similarly, since the claim recites "a nucleotide sequence," which reads on a single base, it is not known how a single base could be used to identify a cell.

For these reasons, the Examiner to hold that undue experimentation is necessary to practice the invention as claimed.

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7. Claim Rejections - 35 USC § 112, first paragraph – written description

A. Claim 1 is rejected under 35 U.S.C. 112, first paragraph, as containing subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention.

This is a genus claim. Nucleic acid molecules which "hybridize" to SEQ ID NO:1 would have one or more nucleic acid substitutions, deletions, insertions and/or additions to said sequence. Similarly, fragments comprising 15 bases of SEQ ID NO:1 would have one or more bases deleted. SEQ ID NO:1 is over 4000 bases and Applicants have not described which of the large number of probes could be used as a marker for the claimed cell, especially in light of the fact that "fragments" as small as 15 bases, or even as small as one base, can be used to specifically identify the claimed cell from other known cell types. Since the claim recites "a nucleotide sequence" a single base reads on the claim. Again, Applicants have not provided sufficient guidance as to how a single base can identify the claimed cell type.

The specification and claims do not indicate what distinguishing attributes are shared by the members of the genus. Thus the scope of the claims includes numerous structural variants, and the genus is highly variant because a significant number of structural differences between genus members is permitted. The specification and claims do not provide any guidance as to what changes should be made. Structural features that could distinguish compounds in the genus from others in the nucleic acid or protein class are missing from the disclosure. No common structural attributes identify the members of the genus. The general knowledge and level of skill in the art do not supplement the omitted description because specific, not general, guidance is what is needed. Since the disclosure fails to describe the common attributes or characteristics that identify members of the genus, and because the genus is highly variant, SEQ ID NO:2, 5 and 32, or molecules which hybridize to the polynucleotides encoding these SEQ ID NOs (which could be at least thousands of molecules) alone are insufficient to describe the genus. One of skill in the art would reasonable conclude that the disclosure fails to provide a representative number of species to describe the genus. Thus, Applicant was not in possession of the claimed genus at the time the invention was made.

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8. Claim Rejections - 35 USC § 112, second paragraph

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

A. Claim 1 is vague and indefinite since claim 1 recites "stringent conditions." It is not known what these conditions are. Nucleic acid molecules which hybridize under conditions of "low" stringency would not necessarily hybridize under conditions of "high" stringency. Furthermore, not all conditions of "high" or "low" stringency, for example, are the same. Therefore, it is required that Applicants amend the claims to recite the exact hybridization conditions without using indefinite phrases such as "for example" without adding new matter.

9. Claim Rejections - 35 USC § 102

A person shall be entitled to a patent unless -

- (b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.
- (e) the invention was described in (1) an application for patent, published under section 122(b), by another filed in the United States before the invention by the applicant for patent or (2) a patent granted on an application for patent by another filed in the United States before the invention by the applicant for patent, except that an international application filed under the treaty defined in section 351(a) shall have the effects for purposes of this subsection of an application filed in the United States only if the international application designated the United States and was published under Article 21(2) of such treaty in the English language.

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A. Claim 1 is rejected under 35 U.S.C. 102(e) as being anticipated by ***. Claims 13-16 of US20060239978.

The applied reference has a common inventor with the instant application. Based upon the earlier effective U.S. filing date of the reference, it constitutes prior art under 35 U.S.C. 102(e). This rejection under 35 U.S.C. 102(e) might be overcome either by a showing under 37 CFR 1.132 that any invention disclosed but not claimed in the reference was derived from the inventor of this application and is thus not the invention "by another," or by an appropriate showing under 37 CFR 1.131.

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B. Claim 1 is rejected under 35 U.S.C. 102(b) as being anticipated by Tomita. The claim recites a marker complementary to SEQ ID NO:1, which hybridizes to SEQ ID NO:1, or which is a fragment of SEQ ID NO:1 as small as 1 base. Tomita teach a polynucleotide 100% identical to SEQ ID NO:1 (Sequence Comparison A). Both are identified as Lrp4 proteins. Therefore, even if Tomita did not disclose the fact that their sequence could be used as a marker, it would be inherent that this polynucleotide, which is 100% identical to that of the claimed invention, could be used as a marker.

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10. Double Patenting

The nonstatutory double patenting rejection is based on a judicially created doctrine grounded in public policy (a policy reflected in the statute) so as to prevent the unjustified or improper timewise extension of the "right to exclude" granted by a patent and to prevent possible harassment by multiple assignees. A nonstatutory obviousness-type double patenting rejection is appropriate where the conflicting claims are not identical, but at least one examined application claim is not patentably distinct from the reference claim(s) because the examined application claim is either anticipated by, or would have been obvious over, the reference claim(s). See, e.g., *In re Berg*, 140 F.3d 1428, 46 USPQ2d 1226 (Fed. Cir. 1998); *In re Goodman*, 11 F.3d 1046, 29 USPQ2d 2010 (Fed. Cir. 1993); *In re Longi*, 759 F.2d 887, 225 USPQ 645 (Fed. Cir. 1985); *In re Van Ornum*, 686 F.2d 937, 214 USPQ 761 (CCPA 1982); *In re Vogel*, 422 F.2d 438, 164 USPQ 619 (CCPA 1970); and *In re Thorington*, 418 F.2d 528, 163 USPQ 644 (CCPA 1969).

A timely filed terminal disclaimer in compliance with 37 CFR 1.321(c) or 1.321(d) may be used to overcome an actual or provisional rejection based on a nonstatutory double patenting ground provided the conflicting application or patent either is shown to be commonly owned with this application, or claims an invention made as a result of activities undertaken within the scope of a joint research agreement.

Effective January 1, 1994, a registered attorney or agent of record may sign a terminal disclaimer. A terminal disclaimer signed by the assignee must fully comply with 37 CFR 3.73(b).

A. Claim 1 is provisionally rejected on the ground of nonstatutory obviousness-type double patenting as being unpatentable over claims 13-16 of copending Application No. US20060239978. Although the conflicting claims are not identical, they are not patentably distinct from each other because the claims of the instant application recite a probe comprising at least 15 bases, or which hybridizes to SEQ ID NO:1. The application also claims methods of using the probe and the cell identified by the probe. The '978 application recites using a large fragment of SEQ ID NO:1 as well as the same methods of using.

This is a <u>provisional</u> obviousness-type double patenting rejection because the conflicting claims have not in fact been patented.

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11. Conclusion

A. No claim is allowable.

Advisory information

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Robert Landsman whose telephone number is (571) 272-0888. The examiner can normally be reached on M-Th 10 AM - 7 PM (eastern).

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Brenda Brumback can be reached on 571-272-0961. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

Robert Landsman, Ph.D. Primary Examiner Art Unit 1647 AB013874 mRNA linear ROD 11-NOV-1998 LOCUS 4864 bp Mus musculus mRNA for Low Density Lipoprotein Receptor Related DEFINITION Protein 4, complete cds. ACCESSION AB013874 AB013874.1 GI:3869144 VERSION Low Density Lipoprotein Receptor Related Protein 4. KEYWORDS SOURCE Mus musculus (house mouse) ORGANISM Mus musculus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Muroidea; Muridae; Murinae; Mus. REFERENCE AUTHORS Tomita, Y., Kim, D.H., Magoori, K., Fujino, T. and Yamamoto, T.T. A novel low-density lipoprotein receptor-related protein with type TITLE II membrane protein-like structure is abundant in heart J. Biochem. 124 (4), 784-789 (1998) JOURNAL PUBMED 9756624 REFERENCE 2 (bases 1 to 4864) AUTHORS Kim, D., Tomita, Y. and Yamamoto, T.T. TITLE Direct Submission Submitted (12-MAY-1998) Dongho Kim, Tohoku University, Gene JOURNAL Research Center; 1-1 Tsutsumidori Amamiya Aoba, Sendai, Miyaqi 981-8555, Japan (E-mail:dondon@biochem.tohoku.ac.jp, Tel:81-22-717-8875, Fax:81-22-717-8877) **FEATURES** Location/Qualifiers 1. .4864 source /organism="Mus musculus" /mol type="mRNA" , /db_xref="taxon:10090" CDS 61. .3402 /codon start=1 /product="Low Density Lipoprotein Receptor Related Protein /protein id="BAA34371.1" /db xref="GI:3869145" /translation="MGRVSFSVRVSSVRRARCSCPGRCYLSCRVPPTTALRALNGLGC AGVPGETAGGAVGPGPLGTRGFLSGSKFQAPGSWKDCFGAPPAPDVLRADRSVGEGCP QKLVTANLLRFLLLVLIPCICALIVLLAILLSFVGTLKRVYFKSNDSEPLVTDGEARV PGVIPVNTVYYENTGAPSLPPSQSTPAWTPRAPSPEDQSHRNTSTCMNITHSQCQILP YHSTLAPLLPIVKNMDMEKFLKFFTYLHRLSCYQHILLFGCSLAFPECVVDGDDRHGL LPCRSFCEAAKEGCESVLGMVNSSWPDSLRCSQFRDHTETNSSVRKSCFSLOOEHGKO SLCGGGESFLCTSGLCVPKKLQCNGYNDCDDWSDEAHCNCSKDLFHCGTGKCLHYSLL CDGYDDCGDPSDEQNCDCNLTKEHRCGDGRCIAAEWVCDGDHDCVDKSDEVNCSCHSO GLVECTSGQCIPSTFQCDGDEDCKDGSDEENCSDSQTPCPEGEQGCFGSSCVESCAGS ${\tt SLCDSDSSLSNCSQCEPITLELCMNLLYNHTHYPNYLGHRTQKEASISWESSLFPALV}$ QTNCYKYLMFFACTILVPKCDVNTGQRIPPCRLLCEHSKERCESVLGIVGLQWPEDTD ${\tt CNQFPEESSDNQTCLLPNEDVEECSPSHFKCRSGRCVLGSRRCDGQADCDDDSDEENC}$

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4864
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ORIGIN

Alignment Scores:

polyA site

 Pred. No.:
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Percent Similarity: 100.0% Conservative: 0
Best Local Similarity: 100.0% Mismatches: 0
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US-10-543-003-3 (1-1113) x AB013874 (1-4864)

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	Qy		IleProSerThrPheGlnCysAspGlyAspGluAspCysLysAspGlySerAspGluGlu 480	
	Db			
	Qy	481	AsnCysSerAspSerGlnThrProCysProGluGlyGluGlnGlyCysPheGlySerSer 500	
	Db	1501		
	Qy	501	CysValGluSerCysAlaGlySerSerLeuCysAspSerAspSerSerLeuSerAsnCys 520	
	Db	1561		
	Qy	521	SerGlnCysGluProIleThrLeuGluLeuCysMetAsnLeuLeuTyrAsnHisThrHis 540	
	Db	1621	AGTCAATGTGAGCCCATCACTTTGGAACTCTGCATGAATTTGCTCTACAACCATACACAT 1680	
	Qу	541	TyrProAsnTyrLeuGlyHisArgThrGlnLysGluAlaSerIleSerTrpGluSerSer 560	
	Db	1681	TATCCAAATTACCTTGGCCACAGAACTCAAAAGGAAGCGTCCATCAGCTGGGAGTCATCC 1740	
	Qу	561	LeuPheProAlaLeuValGlnThrAsnCysTyrLysTyrLeuMetPhePheAlaCysThr 580	
	Db		CTTTTCCCTGCCCTTGTACAAACCAACTGTTACAAATACCTCATGTTTTTCGCTTGCACC 1800	
•	Qy		IleLeuValProLysCysAspValAsnThrGlyGlnArgIleProProCysArgLeuLeu 600	
	Db	1801	ATTTTGGTTCCAAAGTGTGATGTGAATACAGGACACCCATCCCGCCTTGCAGACTCCTG 1860	

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Qу	601	CysGluHisSerLysGluArgCysGluSerValLeuGlyIleValGlyLeuGlnTrpPro	620
Db	1861	TGTGAGCACTCCAAAGAGCGCTGTGAGTCTGTTCTGGGAATCGTTGGCCTGCAGTGGCCT	1920
Qy	621	GluAspThrAspCysAsnGlnPheProGluGluSerSerAspAsnGlnThrCysLeuLeu	640
Db	1921	GAAGACACCGACTGCAATCAATTTCCAGAGGAAAGTTCAGACAATCAAACTTGCCTCCTG	1980
Qy	641	ProAsnGluAspValGluGluCysSerProSerHisPheLysCysArgSerGlyArgCys	660
Db	1981	CCCAATGAAGATGTGGAAGAATGCTCTCCGAGTCACTTCAAATGCCGCTCGGGACGATGC	2040
Qу	661	ValLeuGlySerArgArgCysAspGlyGlnAlaAspCysAspAspAspSerAspGluGlu	680
Db	2041	GTTCTGGGCTCCAGGAGATGTGACGGCCAGGCTGACTGTGACGACGACAGTGACGAGGAG	2100
Qy	681	AsnCysGlyCysLysGluArgAlaLeuTrpGluCysProPheAsnLysGlnCysLeuLys	700
Db	2101	AACTGTGGTTGTAAAGAGAGAGCTCTTTGGGAATGTCCATTTAATAAGCAATGTCTGAAG	2160
Qy	701	HisThrLeuIleCysAspGlyPheProAspCysProAspSerMetAspGluLysAsnCys	720
Db	2161	CATACATTAATCTGCGATGGGTTTCCAGATTGTCCAGACAGTATGGATGAAAAAAACTGC	2220
Qу	721	SerPheCysGlnAspAsnGluLeuGluCysAlaAsnHisGluCysValProArgAspLeu	740
Db	2221	TCATTTTGCCAAGACAATGAGCTGGAATGTGCCAACCATGAGTGTGCCGCGTGACCTT	2280
Qy	741	<pre>TrpCysAspGlyTrpValAspCysSerAspSerSerAspGluTrpGlyCysValThrLeu </pre>	760
Db	2281	TGGTGCGACGGATGGTCGACTGCTCAGACAGTTCTGATGAATGGGGCTGTGTGACCCTC	2340
Qy	761	SerLysAsnGlyAsnSerSerSerLeuLeuThrValHisLysSerAlaLysGluHisHis	780
Db	2341	TCTAAAAATGGGAACTCCTCATTGCTGACTGTTCACAAATCTGCAAAGGAACACCAC	2400
Qy	781	ValCysAlaAspGlyTrpArgGluThrLeuSerGlnLeuAlaCysLysGlnMetGlyLeu	800
Db	2401	GTGTGTGCTGACGGCTGGCGGGAGACGTTGAGTCAGCTGGCCTGCAAGCAGATGGGTTTA	2460
Qу	801	GlyGluProSerValThrLysLeuIleProGlyGlnGluGlyGlnGlnTrpLeuArgLeu	820
Db		GGAGAACCGTCTGTGACCAAGCTGATCCCAGGACAGGAAGGCCAGCAGTGGCTGAGGTTG	
Qу	821	TyrProAsnTrpGluAsnLeuAsnGlySerThrLeuGlnGluLeuLeuValTyrArgHis	840
Db	-	TACCCCAACTGGGAGAATCTCAATGGGAGCACCTTGCAGGAGCTGCTGGTATACAGGCAC	
Qу		SerCysProSerArgSerGluIleSerLeuLeuCysSerLysGlnAspCysGlyArgArg	
Db		TCCTGCCCAAGCAGAAGTGAGATTTCCCTTCTGTGCTCCAAGCAAG	
Qy		ProAlaAlaArgMetAsnLysArgIleLeuGlyGlyArgThrSerArgProGlyArgTrp	
Db		CCTGCTGCCCGAATGAACAAGAGGATCCTTGGGGGGTCGGACTAGTCGTCCTGGGAGGTGG	
Qy		ProTrpGlnCysSerLeuGlnSerGluProSerGlyHisIleCysGlyCysValLeuIle	
Db		${\tt CCGTGGCAGTGCTCTCTGCAGAGTGAACCCAGTGGACATATCTGTGGCTGTTCCTCATT}$	
Qy		AlaLysLysTrpValLeuThrValAlaHisCysPheGluGlyArgGluAspAlaAspVal	
Db	2761	GCCAAGAAGTGGGTCCTGACAGTTGCCCATTGCTTTGAAGGGAAGAAGACGCTGATGTT	2820

Qy 921 TrpLysValValPheGlyIleAsnAsnLeuAspHisProSerGlyPheMetGlnThrArg 940		
Oy	Qу	
Db 2881 TTTGTGAAGACCATCCTGCTACATCCCGTTACAGTCGAGCAGTGGTAGAACTATGATATC 2940 Qy 961 SerValValGluLeuSerAspAsp1leAsnGluThrSerTyrValArgProValCysLeu 980	Db	
Oy 961 SerValValGluLeuSerAspAspIleAsnGluThrSerTyrValArgProValCysLeu 980	Qу	41 PheValLysThrIleLeuLeuHisProArgTyrSerArgAlaValValAspTyrAspIle 960
Db 2941 AGCGTGGAGCTGAGCGATGATATCAATGAGACAAGCTACGTCAGACCTGCTTA 3000 Qy 981 ProSerProGluGluTyrLeuGluProAspThrTyrCysTyrIleThrGlyTrpGlyHis 1000 Db 3001 CCCAGTCCGGAGGAGTATCTAGAACCAGATACGTACTGCTACACAGGCTGGGGCCAC 3060 Qy 1001 MetGlyAsnLysMetProPheLysLeuGlnGluGlyGluValArgIleIleProLeuGlu 1020	Db	81 TTTGTGAAGACCATCCTGCTACATCCCCGTTACAGTCGAGCAGTGGTAGACTATGATATC 2940
Oy 981 ProSerProGluGluTyrLeuGluProAspThrTyrCysTyrlleThrGlyTrpGlyHis 1000	-	
Db 3001 CCCAGTCCGGAGGAGTATCTAGAACCAGATACGTACTGCTACATCACAGGCTGGGCCAC 3060 Oy 1001 MetGlyAsnLysMetProPheLysLeuGlnGluGlyGluValArgIleIleProLeuGlu 1020 Db 3061 ATGGGCAATAAAATGCCCTTTAAGCTGCAGGAGGGAGAGGTCCGCATTATCCCTCTGGAG 3120 Oy 1021 GlnCysGlnSerTyrPheAspMetLysThrIleThrAsnArgMetIleCysAlaGlyTyr 1040 Db 3121 CAGTGCCAGTCCTATTTTGACATGAAGACCATCACCAATCGGATGATCTGTGCTGGCTAT 3180 Oy 1041 GluSerGlyThrValAspSerCysMetGlyAspSerGlyGlyProLeuValCysGluArg 1060 Db 3181 GAGTCTGGCACCGTGGACTCCTGCATGGGAGACAGCGGTGGGCCTCTGGTTTGTGAACGA 3240 Oy 1061 ProGlyGlyGlnTrpThrLeuPheGlyLeuThrSerTrpGlySerValCysPheSerLys 1080 Oy 1081 ValLeuGlyProGlyValTyrSerAsnValSerTyrPheValGlyTrpIleGluArgGln 1100 Oy 1081 ValLeuGlyProGlyValTyrSerAsnValSerTyrPheValGlyTrpIleGluArgGln 1100 Oy 1101 IleTyrIleGlnThrPheLeuGlnLysLysSerGlnGly 1113		
Qy 1001 MetGlyAsnLysMetProPheLysLeuGlnGluGlyGluValArgIleIleProLeuGlu 1020	Qy	
Db 3061 ATGGGCAATAAAATGCCCTTTAAGCTGCAGGAGGAGGAGGTCCGCATTATCCCTCTGGAG 3120 Qy 1021 GlnCysGlnSerTyrPheAspMetLysThrlleThrAsnArgMetlleCysAlaGlyTyr 1040	Db	01 CCCAGTCCGGAGGAGTATCTAGAACCAGATACGTACTGCTACATCACAGGCTGGGGCCAC 3060
Db 3061 ATGGGCAATAAAATGCCCTTTAAGCTGCAGGAGGGAGGGTCCGCATTATCCCTCTGGAG 3120 Qy 1021 GlnCysGlnSerTyrPheAspMetLysThrIleThrAsnArgMetIleCysAlaGlyTyr 1040	Qy	01 MetGlyAsnLysMetProPheLysLeuGlnGluGlyGluValArgIleIleProLeuGlu 1020
Db 3121 CAGTGCCAGTCCTATTTTGACATGAAGACCATCACCAATCGGATGATCTGTGCTGAT 3180 Qy 1041 GluSerGlyThrValAspSerCysMetGlyAspSerGlyGlyProLeuValCysGluArg 1060	Db	61 ATGGGCAATAAAATGCCCTTTAAGCTGCAGGAGGAGGGCCCGCATTATCCCTCTGGAG 3120
Db 3121 CAGTGCCAGTCCTATTTTGACATGAAGACCATCACCAATCGGATGATCTGTGCTGGCTAT 3180 Qy 1041 GluSerGlyThrValAspSerCysMetGlyAspSerGlyGlyProLeuValCysGluArg 1060	Qу	21 GlnCysGlnSerTyrPheAspMetLysThrIleThrAsnArgMetIleCysAlaGlyTyr 1040
Db 3181 GAGTCTGGCACCGTGGACTCCTGCATGGGAGACAGCGGTGGGCCTCTGGTTTGTGAACGA 3240 Qy 1061 ProGlyGlyGlnTrpThrLeuPheGlyLeuThrSerTrpGlySerValCysPheSerLys 1080	Dp,	
Db 3181 GAGTCTGGCACCGTGGACTCCTGCATGGGAGACAGCGGTGGGCCTCTGGTTTGTGAACGA 3240 Qy 1061 ProGlyGlyGlnTrpThrLeuPheGlyLeuThrSerTrpGlySerValCysPheSerLys 1080	Qу	41 GluSerGlyThrValAspSerCysMetGlyAspSerGlyGlyProLeuValCysGluArg 1060
Db 3241 CCCGGAGGACAGTGGACATTATTTGGTTTAACTTCATGGGGGCTCCGTCTGCTTTTCCAAA 3300 Qy 1081 ValLeuGlyProGlyValTyrSerAsnValSerTyrPheValGlyTrpIleGluArgGln 1100	Db	
Db 3241 CCCGGAGGACAGTGGACATTATTTGGTTTAACTTCATGGGGCTCCGTCTGCTTTTCCAAA 3300 Qy 1081 ValLeuGlyProGlyValTyrSerAsnValSerTyrPheValGlyTrpIleGluArgGln 1100	Qy ·	61 ProGlyGlyGlnTrpThrLeuPheGlyLeuThrSerTrpGlySerValCysPheSerLys 1080
Db 3301 GTTCTGGGACCTGGAGTGTACAGCAATGTGTCTTACTTTGTGGGCTGGATTGAAAGACAA 3360 Qy 1101 IleTyrlleGlnThrPheLeuGlnLysLysSerGlnGly 1113	Db	
Db 3301 GTTCTGGGACCTGGAGTGTACAGCAATGTGTCTTACTTTGTGGGCTGGATTGAAAGACAA 3360 Qy 1101 IleTyrlleGlnThrPheLeuGlnLysLysSerGlnGly 1113	Qy	81 ValLeuGlyProGlyValTyrSerAsnValSerTyrPheValGlyTrpIleGluArgGln 1100
	Db	
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